Serial Number: 09/28/7/7  Changed a file from non-ASCII to ASCII  Changed the margins in cases where the sequence text was "wrapped" does to the next line.  Edited a format error in the Current Application Data section, specifically.  Edited the Current Application Data section with the actual current number. The number inputted by the applicant was   the prior application data; or   other    Added the mandatory heading and subheadings for "Current Application Data".  Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer. Changed the spelling of a mandatory field (the headings or subheadings), specifically:  Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted:   non-ASCII "garbage" at the beginning/end of files;   secretary initials/filename at end of files   page numbers throughout text;   other invalid text, such as   secretary initials/filename at end of files   page numbers throughout text;   other invalid text, such as   secretary initials/filename at end of files   secretary i	war.	llit aut get by scarrier (63/ Russ)
Changed the margins in cases where the sequence text was "wrapped" to the next line.    Changed the margins in cases where the sequence text was "wrapped" to the next line.    Edited a format error in the Current Application Data section, specifically.    Edited the Current Application Data section with the actual current number. The number inputted by the applicant was   the prior application data; or   other     Added the mandatory heading and subheadings for "Current Application Data".    Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.     Changed the spelling of a mandatory field (the headings or subheadings), specifically:    Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:    Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:    Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.    Inserted colons after headings/subheadings. Headings edited included:    Deleted:   non-ASCII "garbage" at the beginning/end of files;   secretary initials/filename at end of files   page numbers throughout text;   other invalid text, such as   Inserted mandatory headings, specifically:    Corrected an obvious error in the response, specifically:    Corrected an error in the Number of Sequences field, specifically:    A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.    Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected:	Seria	REQEIVE
Changed the margins in cases where the sequence text was "wrapped" don't to the next tine.  Edited a format error in the Current Application Data section, specifically.  Edited the Current Application Data section with the actual current number. The number inputted by the applicant was   the prior application data; or   other    Added the mandatory heading and subheadings for "Current Application Data".  Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.  Changed the spelling of a mandatory field (the headings or subheadings), specifically:  Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted: non-ASCII garbage" at the beginning/end of files; secretary initials/filename at end of files page numbers throughout text; other invalid text, such as lnserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:  A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.  Deleted anding stop codon in amino acid sequences and adjusted the "(A)Length: field accordingly (error due to a Patentin bug). Sequences corrected:	_ \ 🗆	Changed a file from non-ASCII to ASCII ENTERED
Edited the Current Application Data section with the actual current number. The number inputted by the applicant was   the prior application data; or   other   Added the mandatory heading and subheadings for "Current Application Data".   Edited the "Number of Sequences" (field. The applicant spelled out a number instead of using an integer.   Changed the spelling of a mandatory field (the headings or subheadings), specifically:   Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:   Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:   Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.   Inserted colons after headings/subheadings. Headings edited included:   Deleted extra, invalid, headings used by an applicant, specifically:   Deleted:   non-ASCII "garbage" at the beginning/end of files;   secretary initials/filename at end of files   page numbers throughout text;   other invalid text, such as   Inserted mandatory headings, specifically:   Corrected an obvious error in the response, specifically:   Edited identifiers where upper case is used but lower case is required, or vice versa.   Corrected an error in the Number of Sequences field, specifically:   A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.   Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length: field accordingly (error due to a Patentin bug). Sequences corrected:	$^{\prime}$	Changed the margins in cases where the sequence text was "wrapped" do the next line.
applicant was   the prior application data; or   other   Added the mandatory heading and subheadings for "Current Application Data".		Edited a format error in the Current Application Data section, specifically
Edited the "Number of Sequences" (field. The applicant spelled out a number instead of using an integer.  Changed the spelling of a mandatory field (the headings or subheadings), specifically:  Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of file page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:  A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.  Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected:		
Changed the spelling of a mandatory field (the headings or subheadings), specifically:  Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted: non-ASCII 'garbage' at the beginning/end of files; secretary initials/filename at end of files		Added the mandatory heading and subheadings for "Current Application Data".
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Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted:   non-ASCII *garbage* at the beginning/end of files;   secretary initials/filename at end of file   page numbers throughout text;   other invalid text, such as    Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:  A *Hard Page Break* code was inserted by the applicant. All occurrences had to be deleted.  Deleted ending stop codon in amino acid sequences and adjusted the *(A)Length:* field accordingly (error due to a Patentin bug). Sequences corrected:		Changed the spelling of a mandatory field (the headings or subheadings), specifically:
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applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted:   non-ASCII "garbage" at the beginning/end of files;   secretary initials/filename at end of files   page numbers throughout text;   other invalid text, such as    Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:  A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.  Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patenth bug). Sequences corrected:		Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
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page numbers throughout text;    other invalid text, such as     Inserted mandatory headings, specifically:     Corrected an obvious error in the response, specifically:     Edited identifiers where upper case is used but lower case is required, or vice versa.     Corrected an error in the Number of Sequences field, specifically:     A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.     Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentln bug). Sequences corrected:		Deleted extra, invalid, headings used by an applicant, specifically:
Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:  A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.  Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected:		Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of file; page numbers throughout text; other invalid text, such as
Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:  A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.  Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentln bug). Sequences corrected:		Inserted mandatory headings, specifically:
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A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.  Deleted <i>ending</i> stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected:		Edited identifiers where upper case is used but lower case is required, or vice versa.
Deleted <i>ending</i> stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected:		Corrected an error in the Number of Sequences field, specifically:
due to a Patentin bug). Sequences corrected:		A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
Other: Seg6-corrected Leu Leu to Ite Leu in 2223) Aprosition 15-16		
		Other: Seg6-corrected Leulen to Itelen in 2223) unlongton for position 15-16
translated to the applicant in the first Office		

<sup>\*</sup>Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT s nd a copy of this form.

3/1/95

37

38

39

40

41

42

43 44

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1

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RAW SEQUENCE LISTING PATENT APPLICATION US/09/281,717

This Raw Listing contains the General Information

DATE: 04/24/2000

TIME: 19:33:37

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           Darimont, Beatrice
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           Feng, Weijun
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           Fletterick, Robert J.
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           Kushner, Peter J.
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           Wagner, Richard L.
 7
           West, Brian
           Yamamoto, Keith R.
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10
           COACTIVATOR BINDING
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     <141> CURRENT FILING DATE: 1999-03-30
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DATE: 04/24/2000 PAGE: RAW SEQUENCE LISTING TIME: 19:33:37

PATENT APPLICATION US/09/281,717

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144

DATE: 04/24/2000

TIME: 19:33:37

PAGE: 4 RAW SEQUENCE LISTING

PATENT APPLICATION US/09/281,717

Input Set: I281717.RAW \*

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## RAW SEQUENCE LISTING PATENT APPLICATION US/09/281,717

DATE: 04/24/2000 TIME: 19:33:37

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**Please Note:**Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

# VERIFICATION SUMMARY PATENT APPLICATION US/09/281,717

DATE: 04/24/2000 TIME: 19:33:37

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	-					<b></b> -												- <b></b> - ·		
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Ogihara, N. 8B19

1631 (Rush)

PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/281,717

DATE: 04/24/2000

TIME: 14:02:59

Input Set: I281717.RAW

This Raw Listing contains the General Information Section and up to first 5 pages.

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Corrected Diskette Needed
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            Darimont, Beatrice
 3
            Feng, Weijun
  4
            Fletterick, Robert J.
 5
            Kushner, Peter J.
            Wagner, Richard L.
 7
            West, Brian
 8
            Yamamoto, Keith R.
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 10
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      <141> CURRENT FILING DATE: 1999-03-30
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Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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